

SEQUENCE LISTING

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Hornik, Vered
<110>
       CONFORMATIONALLY CONSTRAINED BACKBONE CYCLIZED SOMATO
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STATIN ANALOGS
       87534-3000
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      09/734,583
< 140 >
       2000-12-13
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      PatentIn version 3.1
<170>
      1
<210>
      14
<211>
       PRT
<212>
       mammalian
<213>
       1
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Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
                                      10
                                                       Ili mar
                 5
<210>
       2
<211>
<212> PRT
<213> Artificial peptide
<220>
<221> DISULFIDE BRIDGE
      (1)..(1)
<222>
       Cys residues at amino acid positions and 6 form a dis
<223>
ulfide birdg
        е
<220>
       MOD_RES
<221>
       (3)..(3)
<222>
        The Trp residue is the D isomer
<223>
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Port.
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< 400>
Cys Phe Trp Lys Thr Cys
      3
<210>
<211>
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<213> Artificial peptide
<220>
<221> MOD_RES
<222> (1)..(1)
<223> N-Methyl
<220>
<221> MOD_RES
<222> (1)..(6)
      cyclo
<223>
<220>
<221> MOD_RES
<222> (3)..(3)
<223> The Trp residue is the D isomer
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Ala Tyr Trp Lys Val Phe
                 5
 1
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        4
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 <212>
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 <222> (1)..(1)
        The Phe residue is a D isomer
 <223>
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<220>
      MOD_RES
<221>
     (8)..(8)
<222>
      The Thr residue ends with CH2OH
<223>
<220>
<221> DISULFIDE BRIDGE
<222> (2)..(2)
<223> A disulfide bride is formed between Cys residues 2 an
d 7
<220>
<221> MOD RES
<222> (2)..(2)
<223> The Trp residue is a D isomer
< 400 > 4
Phe Cys Phe Trp Lys Thr Cys Thr
                5
<210> 5
<211> 7
<212> PRT
<213> Artificial Peptide
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<221> DISULFIDE
<222> (2)..(2)
<223> A Disulfide Bridge is formed between the Cys residues
 at position
        2 and 6
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<220>
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<222> (1)..(1)
<223> The Phe residue is a D isomer
<220>
<221> MOD RES
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BI
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(4)..(4)
<222>
      The Trp residue is a D isomer
<223>
<220>
      MOD RES
<221>
<222> (7)..(7)
<223> The Thr resiude ends with N2H
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Phe Cys Phe Trp Lys Cys Thr
                5
1
      6
<210>
<211> 8
<212> PRT
<213> Artificial peptide
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<222> (1)..(1)
<223> is a gamma amino butyric acid, diamino butyric acid,
Gly, beta-Al
       a, 5-amino pentanoic acid or amino hexanoic acid; Res
idue 1 is bi
       rdged to Residue 8; Residue 1 also begins with a hydr
ogen, or a m
       ono- or di- saccharide attached
<220>
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<222> (2)..(2)
<223> is (D) or (L) Phe or Tyr
<220>
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<222> (3)..(3)
<223> is (D) or (L)-Trp, or (L)-Phe, (D)- or (L)-1Nal or (D
) or (L)-2Na
       1, or Tyr
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<220>
<221>
      MISC FEATURE
<222>
      (4)..(4)
<223>
       is (D) or (L)-Trp
<220>
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<222> (5)..(5)
<223> is (D) or (L)-Lys
<220>
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<222> (6)..(6)
<223> is Thr, Gly, Abu, Ser, Cys, Val, (D) or (L)-Ala, or (
D) - or (L) - A
       la, or Tyr
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<222> (7)..(7)
<223> is (D) or (L)-Phe, or (D)- or (L)-Ala, Nle, or Cys;
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<222> (8)..(8)
<223> is Gly, Val, Leu, (D) or (L)-Phe, or 1Nal or 2Nal; wi
th a termina
       1 carboxy acid, amide or alcohol group.
<400> 6
Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                5
1
       7
<210>
      7
<211>
<212>
      PRT
<213> Artificial Peptide
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<220>
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<222> (1)..(1)
<223> is (D)- or (L)-Phe, or (D)- or (L)-Ala; wherein Resid
ue 1 is brid
      ged to Residue 6 a bridging group composed of 1 to 5
methyl space
      rs connected to an amide, thioether, thioester, or di
sulfide, fol
       lowed by 1 to 5 methyl spacers
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<222> (2)..(2)
<223> is Tyr or (D) - or (L)-Phe
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<223> is (D)- or (L)-Trp, (D)- or (L)-1Nal, or (D)- or (L
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<223> is Thr, Val, Ser, or Cys
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<223> is Gly or (D)- or (L)-Phe
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<223> is Thr, GABA, (D)- or (L)-1Nal, (D)- or (L)-2Nal, or
  (D) - or (L
       )-Phe
<400> 7
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Xaa Xaa Xaa Lys Xaa Xaa Xaa
                5
1
      8
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      Artificial Peptide
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 acids
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<223> is 1Nal, 2Nal, Beta-Asp (Ind), Gly, Tyr, (D)- or (L)
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       ) - or (L) - Phe
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      (3)..(4)
<222>
<223> may be absent, or are independently Gly, Tyr, 1Nal, 2
Nal, Beta-As
       p (Ind), Gly, Tyr, (D)- or (L)-Ala, or (D)- or (L)-P
he
<220>
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 <223> (D) - or (L) -Trp
 <220>
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 <222> (6)..(6)
       (D) - or (L) -Lys
 <223>
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<220>
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<223> is absent or is Gly, Abu, Cys, Thr, Val, (D)- or (L)-
Ala, or (D)
      - or (L)-Phe
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<221> MISC FEATURE
<222> (8)..(8)
<223> is Cys, (D)- or (L)-Ala, or (D)- or (L)-Phe
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<222> (9)..(9)
<223> is absent or is Val, Thr, 1Nal or 2Nal
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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
<210> 9
<211> 7
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<213> Artificial Peptide
<220>
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 <222> (1)..(1)
      (D)- or (L)-Phe, Tyr or (D)- or (L)-Ala; Residue 1 i
 <223>
 s connected
       to Residue 7 by a bridge comprised of 1 to 5 methylen
 e spacers co
       nncected to an amide, thioether, thioester, or disulf
 ide, followe
        d by 1 to 5 methylene spacers
 <220>
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 <222> (2)..(2)
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<223> (D) - or (L) - Phe, Tyr or (D) - or (L) - Ala;
<220>
<221> MISC FEATURE
<222> (3)..(3)
<223> is absent or is (D)- or (L)-Phe, Tyr or (D)- or (L)-
Ala;
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<221> MISC_FEATURE
<222> (4)..(4)
\langle 223 \rangle is (D) - or (L) Tyr
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<223> is (D) - or (L) - Lys
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<222> (6)..(6)
<223> is absent or is Thr, Val, Cys or (D)- or (L)-Ala
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<222> (7)..(7)
<223> is a (D) - or (L) - Phe, Cys, or (D) - or (L) - Ala
< 400>
      9
Xaa Xaa Xaa Xaa Xaa Xaa
                 5
<210> 10
<211> 7
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<213> Artificial Peptide
<220>
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<221> MISC FEATURE
<222> (1)..(1)
<223> is absent or is (D) - or (L) -Phe or Ala; the bride is
connected t
      o Residue 1 or 2 and Residue 6 or 7, wherein the bird
ge is compri
       sed of 1 to 5 methylene spacers connected to an amide
, thioether,
        thioester, or disulfide, followed by 1 to 5 methylen
e spacers
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> is (D) - or (L) - Phe or Ala, Tyr
<220>
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<222> (3)..(3)
<223> is (D)- or (L)-Trp
<220>
<221> MISC FEATURE
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<223> is (D) - or (L) -Lys
<220>
<221> MISC FEATURE
<222> (5)..(5)
<223> is Thr, Ala, Val, or Cys
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       is absent or is (D) - or (L) - Phe, Ala, or Cys
 <223>
 <220>
 <221> MISC_FEATURE
 <222> (7)..(7)
 <223> is absent or is Thr or Thr reduced to an alcholol
```

<400> 10

Xaa Xaa Xaa Xaa Xaa Xaa 1

Blachele